

9 GGCAGGCTGAGGGCAGGTGGGAAGGAAACCCGGGACGCATCGCAGCAGCAGCAGCAGCAGCAGAAGCAGCAGCACGCTGAGCAGCTCGGCAGTCCTTCCAGAGAC	90 CCT TCC CGG CCC CTC CTC CTC TTC TTG CAT CTG GCT TTC CTG GGA GGT CGT TCC CAC CCG CTG GGC Pro Ser Arg Ala Leu Leu Leu Leu Phe Leu Ala Phe Leu Gly Gly Arg Ser His Pro Leu Gly	180 GTG GAG Val. Glu	270 CAC H1s	360 ATC 11e	465 GGC TGC AAA GTG CTG AGG CGG CAT TAAGAGGAAGTCCTGGCTGCAGACACCTGCTTCTGATTCCACAAGGGGCTTTTTCCTCAAACCC Gly Cys Lys Val Leu Arg Arg His	584 12C2
SCO	CTG Leu	GTG Val	666 614	CCC	נעטבנ	10101
Ē	CCG	CAG	CGT	GAC	1110	ATAA
CAGT	614 H138	CTG Leu	ATC Ile	ATG	SCTT	TGAA
1000	700 Ser	GAG	990 914	AAG Lys	שטטט	אָטטט.
, JOSC JOSC JOSC JOSC JOSC JOSC JOSC JOSC	CCT	70G Ser	63.5 <b>61.</b> u	AGG	CACA	110011
מאפט	GGT	CTG Leu	Acc	666 61y	ATTG	AAAT
באטט	663 613	TTG CAG GGC AAA CTG TCG GAG CTG CAG Leu Gln Gly Lys Leu Ser Glu Leu Gln	CTC CNG GNG AGC CCC CGT CCC ACA GGT GTC TGG AAG TCC CGG GAG GTA GCC ACC GNG GGC ATC CGT Leu Gin Glu Ser Pro Arg Pro Thr Gly Val Trp Lys Ser Arg Glu Val Ala Thr Glu Gly Ile Arg	ACC CTG CGG GCA CCA CGA AGC CCC AAG ATG GTG CAA GGG TCT GGC TGC TTT GGG AGG AAG ATG GAC CGG Thr Leu Arg Ala Pro Arg Ser Pro Lys Het Val Gln Gly Ser Gly Cys Phe Gly Arg Lys Het Asp Arg	707	כאכא
ACCA	CTC Leu	660 61y	GTA Val	757 Cys	1661	TGAG
CAGA	TTC Phe	CAG Gln	646 61u	917	כאככ	מכננו
SCAG	GCT	TTG	CGG	TOT	c รอง	CTTA
AGCA(	CTG	TTA CAG GAG CAG GGC AAC CAT Leu Gln Glu Gln Arg Asn His	TCC Ser	666 614	GCTĢ	GTTT
CAGC	CAT	AAC Asn	AAG Lys	CAA Gln	CCTG	GATG
פכאס	TTG	CCC Arg	100 1 r p	GTG Val	AAGT	ataa
CCCA	TTC Phe	CAG	GTC	A TG Het	ดูงดูด	TTAT
CCAT	CTC Leu	GAG	GGT	AAG	TAA	TGTT
CCAC	CTG Leu	CAG	ACA The	000 BIA	CAT HIS	TGAT
ναας	CTC Leu	TTA	000 Pro	AGG	CCC	TATT
GCAA	CTG Leu	666 614	CGT	CGA	AGG	TATT
CGNA	CTC	TOC	ភូពិ ភូពិ	CCA	CTG	TATG
CCTC	GCG	ACG	AGG	GCA	CTG	TATI
GGCA	CGG	GAC TTG GAA ACG TCC GGG ' Asp Leu Glu Thr Ser Gly '	; GAG	CGG	AAA Lys	TAATO
TGAG	700	110	CAG	53	7 160	1111
7660	Pro		CT CT		9 6	77.43
200 000	ACA GCA The Ala	TCG Ser	014	TAC TYF	CTC Leu	3,000
•	ACA	GCC	GAG Glu	CTC	61)	AAGT
	CAG	JCA	CTG	ATG GTC Het Val	. AGT	1110
	000 0100	. GGT	TCC Ser	ATG	TCC AGT GGC	000
	GAT	CCC	ACA	AAA Lys	7CC Sel	584 Tetescocotticaagtosactcattttatttatttatttatttatttatttatttat
	ATG	AGC	CAG	CGC	AGG	1. 1.

593 TTATAAGGT

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FIG. 3-A

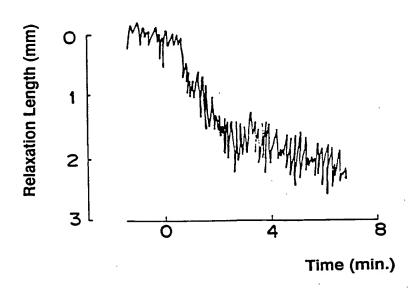


FIG. 3-B

